

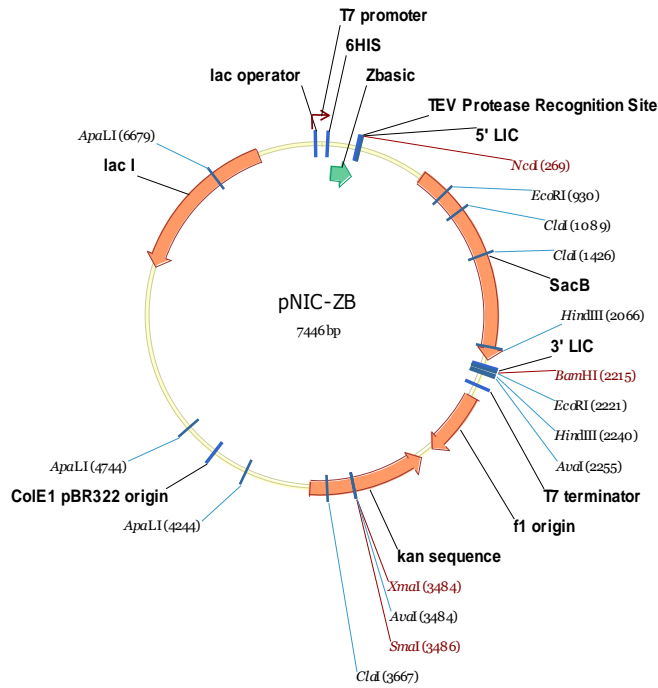
Vector information sheet

Dated: 8th May 2013

Vector Name	pNIC-ZB
Source	Pavel Savitsky
Sequence accession/link	Genbank GU452710

Description	<p>pET expression vector with His₆ and Z-basic (ZB) tags followed by a TEV protease cleavage site.</p> <p>The Z-basic tag (<i>J. Chromatog A</i>, 1161:22-28) is a 54-aa sequence derived from protein A and modified to have a high positive surface charge, allowing the fusion proteins to bind to S-sepharose at salt concentrations in which most cellular proteins do not bind.</p> <p>The vector includes sites for LIC cloning, and a “stuffer” fragment that includes the SacB gene, allowing negative selection on 5% sucrose</p>
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Antibiotic resistance	Kanamycin, 50 µg/ml
Promoter	T7 - lacO
Cloning	LIC. (vector treated with BsaI, then with T4 DNA polymerase in presence of dGTP)
Initiation codon	Supplied in PCR primer
N-terminal fusion – seq.	MHHHHHSSGVDNKFNKERRRARREIRHLPNLNREQRRRAFIR SLRDDPSQSANLLAEAKKLNDAQPKGTEENLYFQ*SM (* - TEV cleavage site)
N-terminal fusion – MW	8975 Da
Termination codons	supplied in PCR primer
Protease cleavage	TEV
Additional features	
Preferred host	DE3 hosts: BL21, Rosetta, etc. MUST express T7 RNA polymerase.
5' sequencing primer	pLIC-for: TGTGAGCGGATAACAATTCC
3' sequencing primer	pLIC-rev: AGCAGCCAACCTCAGCTTCC



Polylinker region:

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6HIS
                                     ~~~~~~
                                     NdeI
                                     ~~~~~~
                                     M H H H H H H H
1  CTAGAAATAA TTTGTTTAA CTTAAGAAG GAGATATACA TATGCACCAT CATCATCATC
   GATCTTTATT AAAACAAATT GAAATCTTC CTCTATATGT ATACGTGGTA GTAGTAGTAG
   6HIS
   Zbasic
   ~~~~~~
61  · S S G V D N K F N K E R R R A R R E I R ·
   ATTTCTTGG TGTGGATAAC AAGTTCAACA AGGAGCGTCG AAGAGCTCGC CGTGAAATTC
   TAAGAAGACC ACACCTATTG TTCAAGTTGT TCCTCGCAGC TTCTCGAGCG GCACTTTAAG
   Zbasic
   ~~~~~~
121 · H L P N L N R E Q R R A F I R S L R D D ·
   GCCATCTGCC GAACCTGAAC CGCGAACAGC GTCGCGCATT TATTTCGAGC CTGCGCGATG
   CGGTAGACGG CTTGGACTTG GCGCTTGTG CAGCGCGTAA ATAAGCGTCG GACGCGCTAC
   Zbasic
   ~~~~~~
181 · P S Q S A N L L A E A K K L N D A Q P K ·
   ATCCGAGCCA GAGCGGAAC CTGCTGGCGG AAGCGAAGAA GCTGAACGAT GCGCAGCCGA
   TAGGCTCGGT CTCGCGCTTG GACGACCGCC TTCGCTTCTT CGACTTGCTA CGCGTCGGCT
   5' LIC
   ~~~~~~
Zbasic TEV Protease Recognition Site
~~~~~
                                     NcoI
                                     ~~~~~~
241 · G T E N L Y F Q S M
   AGGTACCGA GAACCTGTAC TTCCAATCCA TG
   TCCATGGCT CTTGGACATG AAGGTTAGGT AC
   .....(SacB fragment).....
                                     BsaI Lower-LIC BamHI EcoRI SacI
                                     ~~~~~~
2010 GATATCCTAT TGGCATTGAC GGTCTCCAGT AAAGGTGGAT ACGGATCCGA ATTCGAGCTC

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CTATAGGATA ACCGTAACTG CCAGAGGTCA TTTCCACCTA TGCCTAGGCT TAAGCTCGAG

SalI
HindIII
*****~::~~

2070 CGTCGACAAG CTTGCGGCCG CACTCGAGCA CCACCACCAC CACCACTGAG ATCCGGCTGC
GCAGCTGTTC GAACGCCGGC GTGAGCTCGT GGTGGTGGTG GTGGTGACTC TAGGCCGACG
T7-reverse

2130 TAACAAAGCC CGAAAGGAAG CTGAGTTGGC TGCTGCCACC GCTGAGCAAT AACTAGCATA
ATTGTTTCGG GCTTTCCTTC GACTCAACCG ACGACGGTGG CGACTCGTTA TTGATCGTAT
pLIC-rev

Primers for LIC cloning:

Upstream: add TACTTCCAATCCATG to the 5' end (ATG in-frame with the desired coding sequence).

Downstream: add TATCCACCTTTACTG to 5' end of downstream primer; add termination codon, if necessary.

pNIC-ZB sequence:

ctagaataatTTTTgtttaactttaagaaggagatatacatatgcaccatcatcatcatcattcttctg
gtgtggataacaagttcaacaaggagcgtcgaagagctcgccgtgaaattcgccatctgccaacctga
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